OIPE

RAW SEQUENCE LISTING   DATE: 07/05/2001
PATENT APPLICATION: US/09/816,756A   TIME: 13:06:24

Input Set : **A:\SEQ LST - revised Missing Parts response.txt**
Output Set: **N:\CRF3\07032001\I816756A.raw**

Does Not Comply
Corrected Diskette Needed

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 3 <110> APPLICANT: University of Connecticut
 4        Mayer, Bruce
 6 <120> TITLE OF INVENTION: Coiled-Coil Mediated Heterodimerization Functional
Interaction Trap
 8 <130> FILE REFERENCE: 883933.0062
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C--> 10 **<140> CURRENT APPLICATION NUMBER: US/09/816,756A**
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11 <141> CURRENT FILING DATE: 2001-03-24
13 <150> PRIOR APPLICATION NUMBER: 60/141,896
14 <151> PRIOR FILING DATE: 1999-06-30
16 <150> PRIOR APPLICATION NUMBER: PCT/US00/17929
17 <151> PRIOR FILING DATE: 2000-06-29
19 <160> NUMBER OF SEQ ID NOS: 2
21 <170> SOFTWARE: PatentIn version 3.0
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ERRORED SEQUENCES

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40 <210> SEQ ID NO: 2
41 <211> LENGTH: 165
42 <212> TYPE: DNA
43 <213> ORGANISM: Artificial Sequence
45 <220> FEATURE:
47 <223> OTHER INFORMATION: Kozak translation start site, follwed by a Myc-epitope,
followed
48        by a WIN-ZIP-B1 synthetic amphipathic helix, followed by an in-fr
49        ame BamHI cloning site
51 <400> SEQUENCE: 2
52 accatggagc aaaagctcat ttctgaagag gacttgaatg aaggatctac catgtccgtt      60
54 gacgaactgc aggctgaggt tgaccagctg caggacgaga attacgctct gaagaccaag     120
56 gttgcgcagc tgcgtaaaaa ggtggaaaag ctgggaggag gatcc                     165
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E--> 60 c&LdOc:.stmLibl:891220.1 06/12/01

VERIFICATION SUMMARY   DATE: 07/05/2001
PATENT APPLICATION: **US/09/816,756A**   TIME: 13:06:25

Input Set : **A:\SEQ LST - revised Missing Parts response.txt**
Output Set: **N:\CRF3\07032001\I816756A.raw**

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:60 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:184 SEQ:2
L:60 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:24
L:60 M:112 C: (48) String data converted to lower case,
L:60 M:252 E: No. of Seq. differs, <211>LENGTH:Input:165 Found:184 SEQ:2